Axial and Radial Oxylipin Transport^{1[OPEN]}

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Jasmonates are oxygenated lipids (oxylipins) that control defense gene expression in response to cell damage in plants. How mobile are these potent mediators within tissues? Exploiting a series of *13-lipoxygenase (13-lox)* mutants in Arabidopsis (*Arabidopsis thaliana*) that displays impaired jasmonic acid (JA) synthesis in specific cell types and using JA-inducible reporters, we mapped the extent of the transport of endogenous jasmonates across the plant vegetative growth phase. In seedlings, we found that jasmonate (or JA precursors) could translocate axially from wounded shoots to unwounded roots in a LOX2-dependent manner. Grafting experiments with the wild type and JA-deficient mutants confirmed shoot-to-root oxylipin transport. Next, we used rosettes to investigate radial cell-to-cell transport of jasmonates. After finding that the LOX6 protein localized to xylem contact cells was not wound inducible, we used the *lox234* triple mutant to genetically isolate LOX6 as the only JA precursor-producing LOX in the plant. When a leaf of this mutant was wounded, the JA reporter gene was expressed in distal leaves. Leaf sectioning showed that JA reporter expression extended from contact cells throughout the vascular bundle and into extravascular cells, revealing a radial movement of jasmonates. Our results add a crucial element to a growing picture of how the distal wound response is regulated in rosettes, showing that both axial (shoot-to-root) and radial (cell-to-cell) transport of oxylipins plays a major role in the wound response. The strategies developed herein provide unique tools with which to identify intercellular jasmonate transport routes.

Both animals and plants produce potently active lipidderived mediators in response to wounding. These oxylipins (oxygenated lipid derivatives) include leukotrienes and prostaglandins in animals (Funk, 2001) and jasmonates in plants (Wasternack and Hause, 2013). Although these regulators frequently show structural similarities (many are cyclopentenone and cyclopentanone lipids), they operate through different signaling pathways often involving large protein complexes. For example, prostaglandins signal in part through G protein-coupled receptor complexes (Furuyashiki and Narumiya, 2011; Kalinski, 2012), and plant jasmonate signaling operates through the Skp/Cullin/F-box CORONATINE

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INSENSITIVE1 complex (Browse, 2009). Many oxylipins produced in response to tissue damage in metazoans act as paracrine signals to elicit defense responses in distal undamaged cells (Funk, 2001). Similarly, it is possible that jasmonates, including the biologically active derivative jasmonoyl-Ile (JA-Ile; Fonseca et al., 2009), might be transported from cell to cell in plants. However, to date, the majority of studies on oxylipin transport in plants have used exogenous jasmonates, and it remains unclear to what extent these compounds are transported between cells and tissues when produced endogenously.

Based on the fact that jasmonic acid (JA) or methyl jasmonate treatments can affect defense gene expression at a distance to the sites of their application, JA was proposed to operate as a paracrine signal capable of being transported from cell to cell in tomato (Solanum lycopersicum) leaves (Farmer et al., 1992). Similar conclusions were drawn for JA in wild tobacco (Nicotiana sylvestris; Zhang and Baldwin, 1997). Isotope-labeling experiments using exogenous jasmonates have indicated JA/JA-Ile transport away from the site of application to distal tissues and even distal organs (Zhang and Baldwin, 1997; Thorpe et al., 2007; Sato et al., 2011). Additionally, grafting experiments in tomato were consistent with long-distance transport of JA/JA precursors (Li et al., 2002; Schilmiller and Howe, 2005), although other studies did not find evidence for JA transport from wounded leaves to distal unwounded leaves (Strassner et al., 2002). Concerning Arabidopsis (Arabidopsis thaliana), Koo et al. (2009) concluded that

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Jasmonate Transport

JA-Ile accumulation detected in leaves distal to the wound site resulted mainly from de novo synthesis in undamaged leaves rather than from the transport of JA/JA-Ile from the wound site. Recently, a transporter (GLUCOSINOLATE TRANSPORTER1) capable of importing JA-Ile (but not JA) into *Xenopus laevis* oocytes has been described (Saito et al., 2015), further supporting the possibility that jasmonates move between cells.

In addition to the transport of jasmonates, there is much evidence consistent with other wound signaling mechanisms that lead to JA synthesis and JA-mediated defense responses at various distances from wounds. That is, wound-activated signaling pathways can be classified as those working near the damage site (i.e. local responses) and those operating distal to it (Rhodes et al., 2006; Wu et al., 2007). Both these types of wound responses can be difficult to study, because several types of events (including the transport of jasmonates) may contribute to JA signaling. However, there has been some progress in understanding long-distance signaling leading to distal wound responses. These mechanisms include electrical and potentially, hydraulic signaling (for review, see Koo and Howe, 2009; Farmer et al., 2014). Membrane hyperpolarizations have been recorded in wounded plants (Zimmermann et al., 2009); however, their relationship with JA synthesis or JA responses has not yet been reported. In Arabidopsis, wounding of adult-phase rosettes stimulates the leaf-toleaf propagation of signals that (1) can be detected with surface electrodes as cell membrane depolarizations; (2) are propagated from leaf to leaf in a mechanism that requires several clade 3 GLUTAMATE RECEPTOR-LIKE (GLR) genes, including GLR3.3 and GLR3.6; and (3) can induce JA and JA-Ile accumulation in distal unwounded sites (Mousavi et al., 2013). However, even when electrical signals were compromised in both the wounded and distal leaves of a glr3.3 glr3.6 double mutant, JA responses were affected only in the distal leaf; local responses in the damaged leaf itself were similar to the wild type (Mousavi et al., 2013). Therefore, certain clade 3 GLRs operate in rosettestage plants to extend the range of the wound response, and even if these genes are mutated, wounded rosette leaves still produce jasmonates. In summary, both jasmonates made near wounds and jasmonates made far from wounds in response to distal signals might be subject to transport within the plant.

This study focused on the mobility of endogenous jasmonates produced in response to wounding. Here, we ask: how mobile are endogenous jasmonates generated in aboveground tissues in response to wounding? Our analysis was conducted throughout the vegetative phase and included different tissues that ranged from embryonic leaves (cotyledons) and roots to expanded rosette leaves. We investigated whether a *glr3.3 glr3.6* double mutant that reduces leaf-to-leaf signal propagation in the adult phase (Mousavi et al., 2013) could also reduce cotyledon-to-root wound signaling in seedlings. Results from these electrophysiology

experiments then led us to investigate whether JA (or JA precursors) can translocate from wounded cotyledons to roots. To do this, we used two approaches. One was based on mutants in 13-LIPOXYGENASEs (13-LOXs) that are necessary for an early step in the synthesis of the JA precursor oxophytodienoic acid. All four 13-LOXs in Arabidopsis (LOX2, LOX3, LOX4, and LOX6) are known to contribute to JA synthesis in vivo (Chauvin et al., 2013). First, LOX2 is responsible for the synthesis of a large pool of JA in wounded leaves (Bell et al., 1995), and it also produces precursors for the synthesis of arabidopside defense-related metabolites (Glauser et al., 2009). Second, LOX3 and LOX4 act together to produce the JA required for full male fertility (Caldelari et al., 2011). Third, LOX6 produces jasmonates in roots that are first separated from aerial tissues and then wounded (Grebner et al., 2013). We tested the impact of mutations in the different 13-LOXs on root JA signaling after cotyledon wounding. This was followed by grafting experiments between the wild type and the JA-deficient mutant allene oxide synthase (aos; Park et al., 2002) to test whether JA (or JA precursors) could translocate axially from wounded shoots into undamaged roots.

In addition to its role in wounded roots (Grebner et al., 2013), LOX6 has been implicated in long-distance wound signaling in the adult-phase rosette, where it is necessary for most of the rapid distal expression of the JA-responsive gene JASMONATE ZIM-DOMAIN10 (JAZ10) when another leaf is wounded (Chauvin et al., 2013). This and the fact that the LOX6 promoter is active principally in xylem contact cells (Chauvin et al., 2013) provided us with the opportunity to investigate oxylipin transport within leaves. We confirmed the cellular localization of the LOX6 polypeptide with a LOX6-GUS fusion protein. We then used a *lox234* triple mutant expressing a *JAZ10* reporter to test whether jasmonates could be exported from xylem contact cells. These experiments led to unique insights into the transport of jasmonates across different leaf cell layers.

RESULTS

GLR3.3/3.6 Are Not Necessary for Cotyledon-to-Root Jasmonate Signaling in Seedlings

In the adult rosette phase, functional clade 3 *GLR* genes are necessary for efficient leaf-to-leaf signal propagation, leading to JA accumulation in distal unwounded leaves on wounded plants (Mousavi et al., 2013). To test whether the same GLRs are also involved in cotyledon-to-root wound signaling in seedlings, we compared basal and wound-induced levels of the early wound response JA marker *JAZ10* (Acosta et al., 2013) between the wild type and the *glr3.3 glr3.6* double mutant. The double mutant had no significant impact on basal or wound-inducible *JAZ10* levels in aerial or root tissues after cotyledon wounding (Supplemental Fig. S1). To test whether cotyledon wounding or laser stimulation was able to induce surface electrical

potential changes in undamaged cotyledons and roots of stimulated 5-d-old seedlings, we developed a unique electrophysiology protocol (Supplemental Fig. S2A). In agreement with Acosta et al. (2013), puncturing one cotyledon with a needle induced a strong activity of the secretable JAZ10p:GUSPlussec (JGP) reporter in the wounded cotyledon as well as in the root compared with control unwounded plants (Supplemental Fig. S2, B and C). In contrast, the stimulation of cotyledons with a high-energy laser did not induce a visible *JGP* reporter expression (Supplemental Fig. S2D). Moreover, under the experimental conditions tested, we could not detect differences in wound-activated surface potential (WASP) changes between the wild type and the glr3.3 glr3.6 double mutant in either undamaged cotyledons or roots (Supplemental Fig. S2E). However, rapid laser wounding of cotyledons induced robust surface potential changes (depolarizations) in undamaged cotyledons and roots, serving as positive control for the experimental setup (Supplemental Fig. S2F). For needle wounding, the recorded WASP changes for both genotypes were very low (<5 mV) and variable, whereas for laser stimulation, the observed depolarization patterns were uniform (20-30 mV). Nevertheless, amplitudes of surface potential changes were the same between the wild type and the *glr3.3 glr3.6* double mutant for both treatments (Supplemental Fig. S2, G and H). Although surface potential changes could be measured in cotyledons and roots after laser stimulation, their propagation was not dependent on GLR3.3/ 3.6. In addition, wound signaling from cotyledons to roots seemed to work independently of WASPs, because the effects of needle wounding were efficiently transduced to roots to activate JA responses, but they only weakly and unreliably induced WASPs. Therefore, WASP-independent mechanisms must be responsible for efficient JAZ10 induction in roots of aerially wounded plants. Alternative approaches were used to test if JA (or JA precursors) could translocate to undamaged roots.

LOX2 Is Required to Induce Full *JAZ10* Expression in Roots after Cotyledon Wounding

LOX6 is known to produce the bulk of JA and JA-Ile in roots when they are first separated from aerial tissues and then wounded (Grebner et al., 2013). Here, we analyzed the contribution of 13-LOXs to *JAZ10* expression in roots after cotyledon wounding in intact seedlings. The wild type, *lox2*, *lox6*, *lox26* double mutant, and *lox346* triple mutant (retaining only LOX2 activity) were wounded with a needle on one cotyledon, and after 1 h, shoots and roots were separated and analyzed by quantitative real-time (qRT)-PCR. None of the mutants were significantly different from the wild type in basal *JAZ10* levels (Fig. 1A). In contrast, woundinduced *JAZ10* levels were decreased in the aerial tissue of the *lox2* mutant (to approximately 25% of those seen in the wounded wild type), and they were further reduced in the *lox26* double mutant (Fig. 1B). A similar but even more striking pattern was observed for root tissues where *lox2* and *lox26* double mutants reduced postwound *JAZ10* expression to less than 10% of wild-type levels. The *lox6, lox34,* and *lox346* seedlings did not exhibit a compromised wound response in aerial and root tissues (Fig. 1B). In accordance with the qRT-PCR results, secretable *JGP* reporter activity was strongly attenuated in the *lox2* mutant background compared with the wild type after cotyledon wounding (Fig. 1, C and D).

The *lox2* and *lox346* data indicated that a functional LOX2 is necessary and sufficient to ensure near wildtype levels of wound-induced JAZ10 expression in both aerial organs and roots of young seedlings (Fig. 1B). However, LOX2 transcripts have not been detected in unwounded or wounded Arabidopsis roots (Grebner et al., 2013). Consistent with the work by Vellosillo et al. (2007), we observed *LOX2p:GUS* reporter activity only in cotyledons and emerging leaves and not in roots (Fig. 1E). In our experimental conditions, promoter activity for the other three 13-LOX constructs was also not detected in roots of control or cotyledon-wounded seedlings (Supplemental Fig. S3). In contrast, LOX6 transcripts were detected by qRT-PCR in adult roots under basal conditions, and LOX3 and LOX4 transcripts were induced when the roots were first separated from the shoots and then wounded (Grebner et al., 2013). The LOX2p:GUS reporter displayed induction only in the vicinity of a wound (Fig. 1E). Overall, our results suggested that JA and/or JA precursors might be transported from wounded aerial tissues to roots where they activated JAZ10 expression.

Micrografting Reveals Shoot-to-Root Oxylipin Transport

To test the hypothesis that oxylipins made in shoots could be transported to roots, reciprocal grafts were made between the wild type and the JA-deficient mutant *aos*. Both genetic backgrounds contained a *JAZ10p*: GUS reporter to visualize sites of JA signaling. A wildtype scion grafted onto a wild-type rootstock showed weak basal reporter activity and strong activation in both aerial organs and root vasculature after cotyledon and leaf wounding (Fig. 2, A and B). GUS activity encompassed approximately 59% of the wild-type rootstock length, extending for $5 \pm 3 \text{ mm} (n = 13)$ from the grafting site. In contrast, aos/aos grafted seedlings exhibited low basal JAZ10p:GUS levels, which were not further induced by wounding treatments (Fig. 2, C and D). These results validated the JA dependency of the reporter line and our grafting system. In the aos/wild-type graft combination, the JAZ10p:GUS reporter exhibited low basal activity and failed to be induced in the wild-type rootstock after aerial wounding of the aos scion (Fig. 2, E and F). In contrast, although basal reporter activity was low in wild-type/aos grafted seedlings, the reporter was strongly induced after cotyledon or leaf wounding in



Figure 1. LOX2 is required to induce *JAZ10* expression in roots after cotyledon wounding. qRT-PCR of *JAZ10* expression basally (A) and 1 h after cotyledon wounding (B) in 5-d-old aerial organs and roots of *lox2*, *lox34*, *lox6*, *lox26*, and *lox346* mutants. *JAZ10* transcript levels were normalized to those of *UBC21* and are displayed relative to the expression of wild-type controls that are set to one and indicated with dashed lines. Bars represent the means of three biological replicates (\pm sD), each containing a pool of organs from approximately 60 seed-lings. Complete qRT-PCR data are in Supplemental Data Set S1, data set 1. *JGP* reporter activity in control (C) and wounded wild-type (WT) and *lox2* (D) 5-d-old seedlings. Detection of GUS activity was performed 2 h after wounding. Orange asterisks indicate cotyledon wounding sites. Bars = 0.5 mm.

both wild-type scion and JA-deficient *aos* rootstock (Fig. 2, G and H). Similar to wild-type/wild-type wounded seedlings, reporter activity encompassed approximately 68% of the *aos* rootstock length, extending for $6 \pm 2 \text{ mm}$ (n = 10) from the grafting site.

These data provide evidence for axial oxylipin transport from the wounded wild-type scion into the *aos* rootstock, where *JAZ10* promoter activity is consequently activated.

LOX6 and LOX2 Activate *JAZ10* Expression in Wounded and Distal Leaves of Adult Rosette Plants

Severely wounding a leaf causes the generation of multiple signals that are transmitted axially along veins to activate JA synthesis in leaves distal to the wound (Rhodes et al., 2006; Koo and Howe, 2009; Farmer et al., 2014). However, for defenses to be activated fully, radial signals that reach the mesophyll are also likely to occur. Could the transport of jasmonates from veins be a part of these radial signaling events? To test for the export of jasmonates from veins, we conducted an extensive qRT-PCR analysis of JAZ10 expression 1 h after wounding in single, double, triple, and quadruple 13-lox mutants in adult-stage rosettes. Basal JAZ10 levels were consistently lower than the wild type in all of the mutants examined (Fig. 3A). Single 13-lox mutants and the lox34 double mutant did not significantly affect JAZ10 expression in the wounded leaf 8 (L8; Fig. 3B). However and in agreement with Chauvin et al. (2013), the *lox6* mutant reduced *JAZ10* expression in a distal unwounded leaf 13 (L13; Fig. 3C). Lack of LOX2 and LOX6 activity in the lox26 double mutant strongly reduced JAZ10 expression in wounded L8 and totally abolished the marker gene induction in distal L13. Analysis of *lox234* (retaining only LOX6 activity) and lox346 (retaining only LOX2 activity) triple mutants showed that the individual activities of LOX6 or LOX2 could ensure most of the JAZ10 expression in wounded L8 but that LOX6 played the major role in promoting near wild-type JAZ10 expression in distal L13 of a wounded plant. However, the almost abolished JAZ10 induction in distal leaves of lox26 wounded plants and the capability of *lox346* to slightly induce *JAZ10* expression in distal sites (Fig. 3C) suggest that LOX2 also contributes (to a lesser extent than LOX6) to JA synthesis at a distance from wounds in adult-phase plants. As expected, the lox2346 quadruple mutant was severely compromised in JAZ10 induction in wounded L8, and *JAZ10* expression was abolished in distal L13 (Fig. 3, B and C). Taken together, the data indicate that the activity of LOX6 is sufficient to induce near wildtype JAZ10 expression in wounded and distal leaves of adult rosette plants. We therefore focused on LOX6 and using a LOX6p:LOX6-GUS protein fusion reporter, began by identifying the cells in which the LOX6 protein was present.

LOX6-Derived Jasmonates Are Exported from the Vasculature into the Leaf Blade

LOX6p:LOX6-GUS reporter activity was confined to the leaf vasculature of adult rosette plants (Fig. 4A), and



Figure 2. Evidence for axial shoot-to-root transport of jasmonate (or jasmonate precursors). *JAZ10p:GUS* reporter activity in grafted 13-d-old seedlings of the indicated scion/rootstock genotypes: WT/WT grafts (A and B), *aos/aos* grafts (C and D), *aos/*WT grafts (E and F), and WT/*aos* grafts (G and H). One of the two cotyledons was excised before grafting. GUS staining was performed 3 h after wounding aerial organs with a needle at sites indicated by orange asterisks (B, D, F, and H). Note the presence of reporter activity in the rootstock of the wild-type (WT)/*aos* grafted plant. Black arrowheads indicate grafting sites, and red arrowheads indicate reporter activity in the root. Bars = 1 mm.

in agreement with *LOX6* transcript levels (Grebner et al., 2013; Supplemental Fig. S4), the LOX6 protein fusion reporter was also not wound inducible (Fig. 4, B–D). As in the wild type (Chauvin et al., 2013), the *LOX6p:GUS* expression pattern was restricted to the vasculature in the *lox234* triple-mutant background (Supplemental Fig. S5). We previously noted the activity of a transcriptional *LOX6p:GUS* reporter in xylem

contact cells and occasionally, subtrichomal mounds (Chauvin et al., 2013). However, here, we did not observe LOX6-GUS protein expression in subtrichomal mounds in any rosette leaves, including leaf 4 (L4) from control or wounded plants. These LOX6 protein localization findings suggested a strategy to investigate whether LOX6-derived jasmonates could act as part of the radial cell-to-cell wound-signaling component. Specifically, we followed the expression of the JA-responsive JAZ10p:GUS reporter in the leaf lamina of the *lox234* triple mutant, in which the activity of all 13-LOXs, except LOX6, is abolished.

Basal *JAZ10p:GUS* activity in adult wild-type rosettes was low (Fig. 4E) but readily induced in wounded leaf 3 (L3) as well as in distal L4 in both the vasculature and extraveinal regions (Fig. 4, F-H). Wounding of aos rosettes harboring the JAZ10p:GUS reporter did not increase GUS activity with respect to control plants (Supplemental Fig. S6), validating the JA dependency of the reporter. Similar to the wild type, basal *JAZ10p*: GUS activity was very weak in the *lox*234 triple-mutant background (Fig. 4I), but it was significantly induced after wounding L3 in both local and distal leaves (Fig. 4, J and K). The JA signaling reporter was activated in both veins and extraveinal cells in the lox234 mutant retaining only LOX6 activity (Fig. 4L). In addition to mechanical wounding, the JAZ10p:GUS reporter was activated in extraveinal regions in both the wild type and lox234 mutant challenged with larvae of the generalist herbivore Spodoptera littoralis (Supplemental Fig. S7).

Evidence for Cell-to-Cell Oxylipin Transport in Leaves

To determine which cell types responded to JA in a lox234 triple mutant retaining LOX6 activity only in the vasculature, we sectioned L4 transversally from control (unwounded) plants and plants wounded on L3. Activity of the LOX6p:LOX6-GUS fusion reporter remained confined to the xylem pole, was predominantly in contact cells, was not wound inducible, and did not translocate to other vascular cells in response to wounding (Fig. 5, A and B). Next, we examined the activity of the JAZ10p:GUS reporter. Basal JAZ10p:GUS expression was undetectable in cells of the epidermis, mesophyll, and vascular bundles in both the wild-type and *lox234* backgrounds. However, sections from wounded plants showed robust reporter activation extending from xylem contact cells into cells of the whole vasculature, the mesophyll, and the epidermis in both genetic backgrounds (Fig. 5, C-F; Supplemental Fig. S8). Reporter activity extended $125 \pm 15 \,\mu\text{m}$ (*n* = 10) from xylem contact cells to the abaxial leaf epidermis in the wild type and 132 \pm 22 μ m (n = 10) in the lox234 triple mutant.

DISCUSSION

How do signals that originate in spatially restricted wounds lead to responses elsewhere in the plant body?



Figure 3. LOX6 activity is sufficient to activate near wild-type *JAZ10* expression in wounded and distal leaves of adult rosette plants. qRT-PCR of *JAZ10* expression in unwounded L8 (A) and 1 h after wounding L8 (B) and distal L13 (C) in indicated genotypes. Basal *JAZ10* levels are similar between L8 and L13 (Mousavi et al., 2013); thus, L8 was used as the unwounded control. Note that, after wounding, the *lox234* mutant harboring only LOX6 activity reaches near wild-type *JAZ10* expression levels in both L8 and L13. *JAZ10* transcript levels were normalized to those of *UBC21* and are displayed relative to the expression of wild-type control (L8) or wounded (L8 or L13) levels that are set to one and indicated with dashed lines. Bars represent the means of three to four biological replicates (\pm sD) from individual 4.5-week-old plants. Complete qRT-PCR data are in Supplemental Data Set S1, data set 2.

Key to resolving this longstanding question will be to decipher both axial and radial signaling mechanisms. Here, and bearing in mind that mechanistic aspects and cell types involved in long-distance wound signaling may differ during plant development, we set the goal of characterizing the mobility of endogenous woundinduced jasmonates.

Axial Translocation of JA (or JA Precursors) from Shoots to Roots in Seedlings

Wounding of a single Arabidopsis cotyledon with a needle causes a strong jasmonate response in roots (Acosta et al., 2013; Gasperini et al., 2015; Larrieu et al., 2015). Having found that several clade 3 *GLR* genes

(GLR3.1, GLR3.2, GLR3.3, and GLR3.6) play a role in propagating membrane depolarizations in the adultphase rosette (Mousavi et al., 2013), we expected to find the same mechanism in wounded seedlings. However, our results suggest that two GLRs (GLR3.3 and GLR3.6) that contribute to the propagation of wound signals in adult-phase leaves are not required for cotyledon-to-root wound signaling that leads to root JA responses. Moreover, in seedlings wounded on one cotyledon, we were able to break the correlation between electrical events measured as surface potentials and JA responses in roots. These observations agree with the work by Rhodes et al. (2006), which concluded that (1) readily detectable electrical signals are not produced from small wounds and that (2) they are not necessary for defense signaling over short distances. Other mechanisms, therefore, contribute to activating JA responses in the roots of seedlings when aerial tissues are wounded.

Our studies of root *JAZ10* expression from aerially wounded plants in a variety of *13-lox* mutants suggested that cotyledon-expressed LOXs (chiefly LOX2)



Figure 4. LOX6-derived oxylipins are exported from the vasculature into the leaf blade. A to D, *LOX6p:LOX6-GUS* protein fusion activity in rosettes before (A) and after (B–D) wounding. E to L, *JAZ10p:GUS* activity in the wild type (WT) and *lox234* triple mutant before (E and I) and after (F–H and J–L) wounding. In each case, L3 of 21-d-old plants was wounded (orange asterisks), and GUS staining was performed 6 h later. L4 from a wounded (W) plant is shown in C, G, and K, and details of the primary order (1°), secondary order (2°), and tertiary order (3°) veins in L4 are shown in D, H, and L. Bars = 0.2 cm (A, B, E, F, I, and J), 1 mm (C, G, and K), and 200 μ m (D, H, and L).



Figure 5. Oxylipins produced in xylem contact cells activate *JAZ10* expression throughout the vascular cylinder and in extraveinal cells. A and B, Cross sections through the midvein showing the localization of LOX6-GUS fusion protein expressed under the *LOX6* promoter before (A) and after (B) wounding. C and D, *JAZ10p:GUS* activity in the wild type (WT) before (C) and after (D) wounding. E and F, *JAZ10p:GUS* activity in the *lox234* triple mutant before (E) and after (F) wounding. L4 of 21-d-old plants was stained for GUS activity in control samples and 6 h after wounding L3. c, Xylem contact cell; m, mesophyll; p, phloem region including phloem parenchyma; v, xylem vessel. Bar = 50 μ m.

can produce JA (or JA precursors) that could be then translocated to roots. This finding is in agreement with the work by Zhang and Baldwin (1997), which found that radiolabeled JA applied to wild tobacco leaves can be transported from aerial tissues to roots. Our grafting experiments established that this was also the case for endogenous JA (or JA precursors). Grafting wild-type scions onto *aos* JA biosynthesis mutant rootstocks revealed that, similar to a wild-type/wild-type graft, JA and/or its precursors can be translocated at least 5 mm from the graft junction into the primary root. In doing so, these translocated oxylipins must pass axially through many cells. Phloem is known to be involved in the transport of several plant hormones (Ham and Lucas, 2014). *JAZ10p:GUS* expression was visible along the root vascular bundle, suggesting that shoot-to-root oxylipin transport might use the phloem route.

In wild tobacco, Zhang and Baldwin (1997) found that JA pools in roots increased 90 min after damage to aerial tissues. The kinetics of the major phase of JA accumulation signaling in Arabidopsis roots after the injury of aerial tissues are similar. In Arabidopsis, shoot-to-root wound signals generated in response to a puncture wound on one cotyledon caused an initial rapid but minor wave of root JA signaling that takes place in the first 20 min after damage (Larrieu et al., 2015). Subsequently, after an apparent delay lasting approximately 20 min, a second major wave representing approximately 85% of total jasmonate signaling output occurs in the root between 40 and 120 min after cotyledon wounding (Larrieu et al., 2015). Because grafting experiments revealed a very strong contribution of shoot-derived JA (or JA precursors) to root JA signaling, we speculate that jasmonate translocation may correlate with the second major burst of JA signaling observed by Larrieu et al. (2015). That is, the axial transport of jasmonates may be the predominant mechanism of shootto-root JA signaling at the seedling stage. This leaves open the possibility of an initial and more rapid shootto-root signaling event that is not necessarily explained by the translocation of JA (or JA precursors) that we observed. This will require further attention. As stated in Rhodes et al., 2006, there is perhaps no need for electrical signals (i.e. transmembrane ion fluxes that can be detected with surface electrodes) to activate JA responses within short distances (here defined as <1 cm) of a wound (1 cm corresponds to the length of an entire 5-d-old Arabidopsis seedling). Finding evidence of oxylipin transport in seedlings led us to test whether this also occurred in leaves.

Radial Transport of Jasmonates from Veins into Interveinal Leaf Regions

Organ-to-organ wound signaling has been studied extensively in adult-phase plants. To date, much of the literature has implicated two tissues as principal routes for long-distance wound signals: phloem and xylem (for review, see Koo and Howe, 2009; Farmer et al., 2014; van Bel et al., 2014). The phloem was long ago identified as an excitable tissue that propagates damage signals through Mimosa pudica (Bose, 1926). Recent evidence from a variety of other plants supports this (for review, see van Bel et al., 2014) along with real-time recordings of wound-induced electrical events in Arabidopsis sieve cells (Salvador-Recatalà et al., 2014). Moreover, sites of JA synthesis in tomato are associated with the phloem (Hause et al., 2003). In parallel, the xylem may also play a crucial role in axial wound signal transmission by allowing the propagation of pressure waves (Stahlberg and Cosgrove, 1997; Farmer et al., 2014), and sites of JA synthesis are present adjacent to xylem vessels in Arabidopsis (Chauvin et al., 2013). Although LOX6 contributes to JA production in the vicinity of the wound, it has a dominant impact on promoting JA/JA-Ile synthesis at a distance from wounds in distal undamaged leaves (Chauvin et al., 2013). It is possible that, during leaf-to-leaf wound signaling, some jasmonates produced by LOX6 in xylem contact cells are then transported into phloem sieve elements or xylem vessels, where they could be further translocated. Such scenarios remain to be tested.

By using an antibody-based approach, Mielke et al. (2011) found that jasmonates are distributed within all cells near the site of damage of mechanically wounded tomato leaves. Herein, we showed that wounding a leaf of the wild type or the *lox234* triple mutant harboring LOX6 activity in xylem contact cells induced JAZ10 expression in cells outside the region of LOX6 protein accumulation. In fact, we were able to detect GUS staining from the wound-activated expression of JAZ10p:GUS in a region encompassing at least six cell types other than contact cells: xylem parenchyma, vascular cambium, cells in the phloem pole, bundle sheath cells, mesophyll cells, and epidermis. LOX6-derived jasmonates are, therefore, transported radially (outward) from xylem contact cells. The distance from xylem contact cells to the abaxial epidermis was about 130 μ m. Under our experimental conditions, it is evident that contact cell-produced oxylipins can be transported radially over at least eight cell layers (Supplemental Fig. S8).

The *lox234* mutant (retaining only LOX6 activity) also expressed near wild-type levels of the JAZ10p:GUS reporter in leaves challenged with S. littoralis. Both the wild type and the *lox234* triple mutant showed strong reporter activation in the center of insect-fed rosettes (which are often the most defended tissues in insect feeding experiments), including in plants that express only LOX6 (Chauvin et al., 2013). However, the early wound response marker gene JAZ10 is not always a suitable marker for defense responses (Gasperini et al., 2015), and it is not surprising that the lox 234 mutant is more susceptible than the wild type to herbivorous insects (Chauvin et al., 2013). In fact, the lox234 mutant lacks LOX2 activity, which is known to contribute to leaf defense against S. littoralis (Glauser et al., 2009). Additionally, the transport of jasmonates from LOX6expressing cells may not be sufficient to cover areas of the leaf distant from the site of damage.

Which Oxylipins Are Mobile in Wounded Arabidopsis?

After mechanical damage, JA and JA-Ile are abundant oxylipins in many leaf cell types (Mielke et al., 2011), and both compounds are possible candidates for mobile jasmonates as suggested previously (Farmer et al., 1992, 2014; Zhang and Baldwin, 1997; Li et al., 2002; Schilmiller and Howe, 2005; Thorpe et al., 2007; Wu et al., 2007; Mielke et al., 2011; Sato et al., 2011). In Arabidopsis seedlings, the gene encoding the JASMONIC ACID RESISTANT1 (JAR1) enzyme (Staswick and Tiryaki, 2004) that converts JA to JA-Ile was found to be indispensable for the activation JA signaling in roots after wounding cotyledons, although JA signaling occurred in the wounded aerial organs of the jar1-1 mutant (Acosta et al., 2013). This evidence makes JA-Ile an unlikely candidate for a mobile jasmonate in shoot-to-root translocation under our experimental conditions. For Arabidopsis leaves, it is at present not possible to conclude which forms of jasmonate are transported, although oxophytodienoic acid was excluded from leaf-to-leaf transport in experiments by Koo et al. (2009). Interestingly, lesion-bearing leaves of the accelerated death2 mutant (Greenberg et al., 1994), in which oxylipin synthesis is hyperactivated, accumulate JA and to a lesser extent, its intermediate precursor 3-oxo-2-(2-(Z)-pentenyl)cyclopentane-1-butanoic acid (OPC4; Mueller et al., 2006). In the same study, JA predominates over OPC intermediates in wounded wild-type Arabidopsis plants. JA and 3-oxo-2-(2-(Z)-pentenyl)-cyclopentane-1butanoic acid are considered here as the most likely transported oxylipins; however, this hypothesis will require careful verification.

CONCLUSION

Our current understanding of oxylipin transport is still rudimentary, but herein, we show that the transport of endogenous wound-induced jasmonates can occur in two vectors relative to organ growth: axial and radial. In the timeframe tested, we present evidence that JA (or JA precursors) can travel a minimum of 0.5 cm from wounded shoots to undamaged roots. This and the finding of radial oxylipin transport in leaves help to explain why the treatment of plants with exogenous jasmonates is so effective in stimulating defense gene expression in distal untreated tissues. This work shows an interesting parallel to prostanoid transport in metazoans.

MATERIALS AND METHODS

Plant Material and Growth Conditions

Arabidopsis (*Arabidopsis thaliana*) accession Columbia-0 was the wild type and the background for previously described mutants and reporter lines used in this study: *glr3.3 glr3.6 (glr3.3a glr3.6a*; SALK_099757 SALK_091801; Mousavi et al., 2013), *JGP* reporter line (Acosta et al., 2013), *aos* (Park et al., 2002), *lox2* (*lox2-1*; Glauser et al., 2009), *lox3* (*lox3B*; SALK_147830), *lox4* (*lox4A*; SALK_071732), *lox6* (*lox6A*; SALK_138907), *lox26*, *lox34*, *lox2346*, and *lox2346* (Caldelari et al., 2011; Chauvin et al., 2013). After seed stratification for 2 d at 4°C, plants were grown at 21°C under 100 μ E m⁻² s⁻¹ light, with a photoperiod depending on the application (seedlings, 14-h-light/10-h-dark cycle; rosette plants, 10-h-light/14-h-dark cycle). For seedling growth, seeds were surfaced sterilized and grown on one-half-strength solid Murashige and Skoog medium (MS; 2.15 g L⁻¹, pH 5.7; Duchefa) supplemented with 0.5 g L⁻¹ MES hydrate (Sigma) and 0.7% (w/v) agar (for horizontally grown seedlings) or 0.85% (w/v) agar (for vertically grown seedlings) as described (Acosta et al., 2013).

Plant Treatments

Wounding of 5-d-old and grafted seedlings was performed with a 25-gauge \times 5/8-in needle (0.5 \times 16 mm) under a stereomicroscope. For wounding of adult rosettes, leaves were numbered from oldest to youngest, and 50% of selected

leaves were crushed with metal forceps in distal parts from the petiole. For insect feeding experiments, three neonate *Spodoptera littoralis* larvae were placed on individually grown 21-d-old plants located in Plexiglas boxes for 24 h before GUS staining.

Electrophysiology

Seedlings were grown vertically in petri dishes on sterile 11- \times 11-cm, 3-mm CHR Whatman Paper (GE Healthcare) placed on solid medium (one-halfstrength MS, 0.5 g L⁻¹ MES hydrate, pH 5.7, and 0.85% [w/v] agar) for 4 d; 1 d before the experiment, short root portions were gently lifted with a sterile toothpick from the paper without disturbing the hypocotyl or root tip that remained in contact with the moist paper support. Ag/AgCl electrodes were used for surface potential measurements on 5-d-old seedlings. Electrodes were prepared as follows: 4-mm-long pieces of silver wire (0.25 mm in diameter; World Precision Instruments) serving as electrode tips were soldered to 10-mmlong spiral copper wires (0.1 mm in diameter) that were twisted into spirals to serve as movement absorbers. The copper spirals were in turn soldered onto 0.7-mm-diameter copper wires acting as holders between the electrode tip and the amplifier probes. Soldered connections between silver and copper wires were isolated with Parafilm, and the nonisolated part of the silver wire was chloridized by electrolysis in 0.1 M HCl. Contacts between plant surfaces and measurement electrodes were enhanced by the addition of approximately $1 \mu L$ of 10 mM KCl in 50% (v/v) glycerol. Ag/AgCl reference electrodes (0.5 mm in diameter) were placed in the side of the petri dish in direct contact with the growth medium. Contact between the chloridized silver wire and the growth medium was provided by an agar bridge composed of a 6-mm-wide silicon tube half filled with 1 M KCL in 1% (w/v) agar in contact with the medium. The upper part of the silicon tube was filled with aqueous 1 M KCl in which the chloridized silver wire was immersed. Electrophysiological measurements were conducted at a stable temperature of 22°C and light intensity of 55 µmol m⁻² s⁻ photosynthetically active radiation. Measurement electrodes were mounted on high-resistance probes on a two-channel amplifier FD223a (World Precision Instruments), which together with InstruTECH LIH 8+8 Acquisition Interface (HEKA Electronic) and Chartmaster software (Heka Electronic), was used to record surface potentials at a 100-Hz sampling frequency. Laser wounding used an IR (808-nm) light-emitting diode (LED) laser system (MDL-N-808-W LED laser head coupled with the PSU-H-LED Power Supply; Changchun New Industries Optoelectronics Technology Co.) controlled by the InstruTECH LIH 8+8 Acquisition Interface; 0.5-s impulses of light with maximum laser power (approximately 8 W) were supplied to one cotyledon using 0.4-mm-diameter fiber optics (Changchun New Industries Optoelectronics Technology Co.). This punctured a hole in the cotyledon in less than 1 s. Seedling wounding with a needle was performed as above. Data analysis used the Chartmaster software. Amplitudes of depolarizations were automatically calculated with a homemade protocol as the difference between the depolarization minimum recorded in the 200 s after wounding and the average surface potential in the 10 s before wounding.

Grafting

Grafts were performed as in Melnyk et al. (2015) with minor modifications. Vertically grown 5-d-old seedlings were grafted under sterile conditions using a dissecting microscope. Seedlings were transferred to square petri dishes containing one layer of 2.5- × 11-cm sterilized Hybond C Super Nitrocellulose Membrane (RPN.203G; Amersham) on top of sterile 11- × 11-cm, 3-mm CHR Whatman Paper (GE Healthcare) placed on solid medium (one-half-strength MS, 0.5 g L⁻¹ MES hydrate, pH 5.7, and 1.5% agar) to guarantee adequate humidity and nourishment. One cotyledon was removed to facilitate handling, and a transverse cut was made through the hypocotyl as close as possible to the shoot using a stainless steel 23 surgical blade (Heinz Herenz Medizianlbedarf GMBH). Grafts were assembled by butt alignment of the two cut halves (Turnbull et al., 2002). After grafting, petri dishes were sealed with micropore tape and kept horizontal for 8 d. Successful graft formation was evaluated by the attachment of scion to rootstock and the resumption of root growth.

Gene Expression

Root and shoot samples of 5-d-old seedlings were collected as described (Acosta et al., 2013), and leaves from adult rosettes were sampled as in Chauvin et al. (2013). RNA and complementary DNA were prepared as in Gfeller et al.

(2011). qRT-PCR was performed as described (Chauvin et al., 2013). Primers for qRT-PCR of *JAZ10* (At5g13220) and *UBIQUITIN-CONJUGATING ENZYME21* (*UBC21*; At5g25760) have been previously reported in Gfeller et al. (2011). To quantify *LOX6* transcripts (At1g67560), the following primers were used: gaagaaatcatagctcgagaagttg and aatggcaagagtagtagt.

Transgenic Lines

Promoters were amplified from wild-type genomic DNA with indicated oligonucleotides for LOX2 (cggggtacctgacaacaaaattcccgaaa and ttccccccgggggcttacattcctctcatatcca; 2,904 bp; At3g45140), LOX3 (cggggtacccttgatacgcgataatcttacaagc and ttcccccccgggttagtgaagaagaaggagttgagga; 2,875 bp; At1g17420), and LOX4 (cggggtaccgaggccaaggacactaggaa and ttccccccgggcaagactgagtttgagagctttctt; 2,719 bp; At1g72520) and cloned by restriction with XmaI and KpnI into a modified pUC57 (Chauvin et al., 2013) to create pEN-L4-promoter-R1 clones. Underlined sequences represent XmaI and KpnI sites. Fusions to GUS were obtained by Gateway Cloning Technology as described (Chauvin et al., 2013). Promoters for LOX6 (Chauvin et al., 2013) and JAZ10 (Acosta et al., 2013) were described previously. A unique nonsecretable version of the JAZ10p: GusPlus (JAZ10p:GUS) reporter was generated in wild-type, aos, and lox234 backgrounds and used in all experiments described in Figures 2, 4, and 5 and Supplemental Figures S1 to S8. The coding DNA sequence of LOX6 was amplified from wild-type complementary DNA with (atgttcgtagcatctccggtaa and aatggaaatgctgttgggaata) oligonucleotides containing the appropriate recombination adaptors and cloned into pDONR221 (Invitrogen). The LOX6p:LOX6-GUS protein fusion construct was generated by Multisite Gateway Technology of entry clones into pH7m34gw. All constructs were introduced into Arabidopsis backgrounds by floral dip Agrobacterium tumefaciens-mediated transformation (Clough and Bent, 1998). For promoter fusions, transformed seeds expressing red fluorescence protein in T1, T2, and T3 lines were selected by fluorescence microscopy, whereas for the protein fusion reporter, lines were selected on medium containing 40 mg L^{-1} hygromycin. A minimum of two independent transgenic lines was used for each construct to perform experiments and verify reproducibility.

GUS Staining, Sectioning, and Light Microscopy

For GUS staining, samples were prefixed for a minimum of 30 min in 90% acetone, washed two times with 50 mM sodium phosphate buffer (pH 7.2), vacuum infiltrated in the staining solution [10 mM Na₂EDTA, 50 mM sodium phosphate buffer, pH 7.2, 3 mM K₄Fe(CN)₆, 3 mM K₃Fe(CN)₆, 0.1% Triton X-100, and 0.5 mg mL⁻¹ 5-bromo-4-chloro-3-indolyl- β -glucuronic acid], and incubated at 37°C in the dark for 2 h (5-d-old seedlings), 4 h (grafted plants), or overnight (adult rosettes). Five-day-old seedlings and grafted plants were cleared in 70% ethanol, mounted in chloral hydrate:glycerol:water solution (8:2:1), and imaged with a Leica MZ16A Stereomicroscope fitted with a DFC310FX Camera. Adult rosettes were cleared in 70% ethanol and imaged with a Keyence Digital Microscope VHX-500F. Transversal sections of leaves were performed as described (Chauvin et al., 2013), mounted in 10% glycerol, and imaged with a Leica DM5500 Microscope fitted with a DFC420 Camera.

Supplemental Data

The following supplemental materials are available.

- Supplemental Figure S1. qRT-PCR of *JAZ10* expression 1 h after cotyledon wounding in aerial organs and roots of wild-type and *glr3.3 glr3.6* 5-d-old seedlings.
- Supplemental Figure S2. Surface potentials recorded from cotyledons and roots of stimulated seedlings.
- Supplemental Figure S3. LOX3p:GUS, LOX4p:GUS, and LOX6p:GUS expression in control and 2 h after cotyledon wounding wild-type 5-d-old seedlings.
- Supplemental Figure S4. LOX6 transcripts are not wound inducible.
- Supplemental Figure S5. Basal (control) and 6 h after wounding LOX6p: GUS expression in 3-week-old rosettes of the *lox234* triple mutant.
- **Supplemental Figure S6.** Basal (control) and 6 h after wounding *JAZ10p*: *GUS* expression in 3-week-old rosettes of the JA-deficient mutant *aos*.

- **Supplemental Figure S7**. *JAZ10p:GUS* expression in 21-d-old rosettes of the wild type and the *lox234* triple mutant in control conditions or after 24 h of treatment with neonate *S. littoralis* larvae.
- **Supplemental Figure S8.** Oxylipins produced in xylem contact cells activate *JAZ10* expression throughout the vascular cylinder and into the mesophyll and epidermis.

Supplemental Data Set S1. Complete qRT-PCR data.

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